

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSXLSIGDRRLGAALLDQGYLDNQLQRALERHSEVGGRLAELVLYDSGYSERLISXIESEFGIPLVNLHSEVETSPKXK	0
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1104731 ACNS8M RS01410 WP 168417583.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

logo

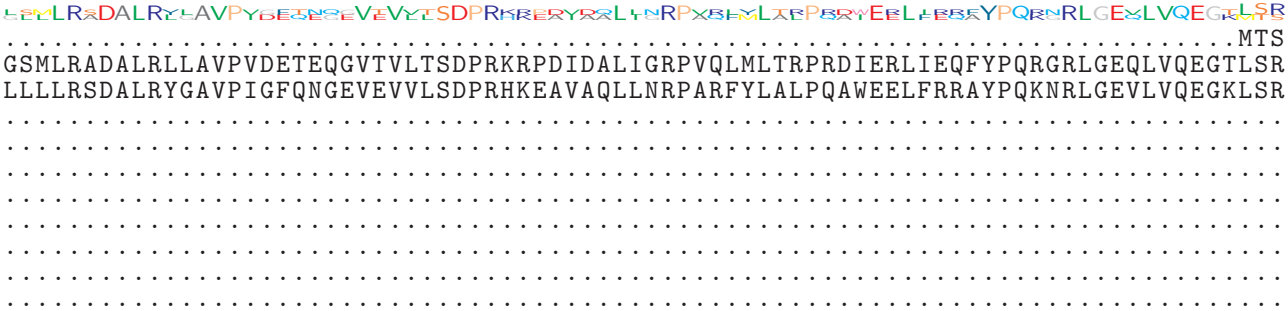
NTDB id 1410 SGL RS13950 WP 010873519.1	SETPAEKAKELRAEPFALDEEADYRVAEYNPLDSESEYEDSGHYEPYQTRSEYMAAATHYPELGLIENYPSGK	0
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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	SBAGSSEKLGEELEYSGYIISSELYEALDQERTGBALGRTLYSSRATPESELYEXLAEEQEGTEFLESFEGMPSEEXK	0
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1
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consensus



3
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logo

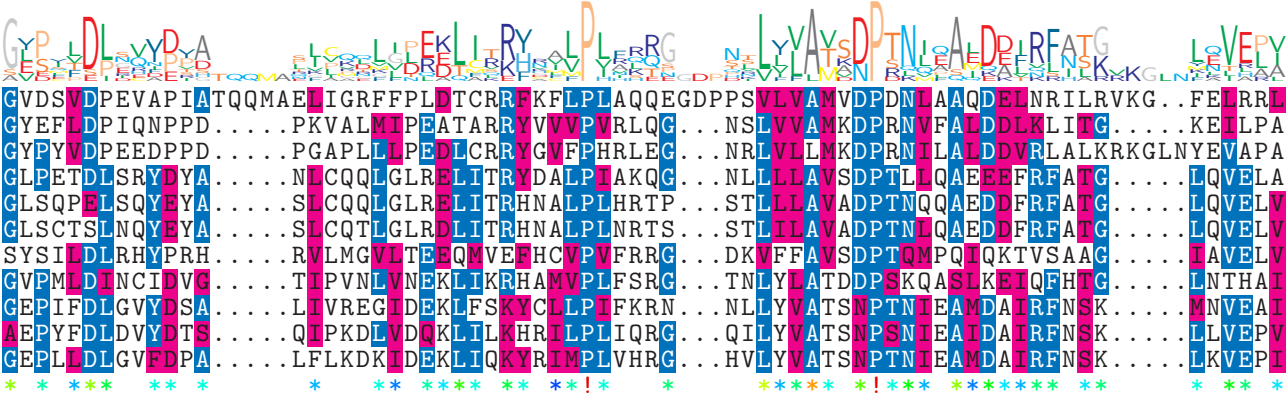
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consensus



83
388
388
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65

logo

NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1306 DR RS10060 WP 027479822.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
NTDB id 1293 VP RS12240 WP 005479695.1
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NTDB id 1110 NGFG RS09235 WP 003689817.1
NTDB id 1251 GC085 RS07725 WP 011946523.1
NTDB id 1104731 ACNS8M RS01410 WP 168417583.1
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consensus



161
455
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124
124
123
132
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132

logo

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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
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consensus		

- ⓧ non conserved
- ✖ similar
- ⓧ ≥ 50% conserved